

# Jorgensen

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## Daten von Jorgensen

```
#Datensortierung

# In Monate umrechnen
jorgensen_r$V1 <- jorgensen_r$V1 * 12
# In Mortalität umrechnen
jorgensen_r$V2 <- 100 - jorgensen_r$V2

jorgensen_r <- jorgensen_r[complete.cases(jorgensen_r),]

jorgensen_r <- jorgensen_r[order(jorgensen_r$V2), ]

# Werte außerhalb des Bereichs [0, inf] raus
jorgensen_r <- jorgensen_r[jorgensen_r$V1 >= 0, ]

xr_jorgensen <- sort(jorgensen_r$V1)
yr_jorgensen <- jorgensen_r$V2

# In Monate umrechnen
jorgensen_b$V1 <- jorgensen_b$V1 * 12
# In Mortalität umrechnen
jorgensen_b$V2 <- 100 - jorgensen_b$V2

jorgensen_b <- jorgensen_b[complete.cases(jorgensen_b),]

jorgensen_b <- jorgensen_b[order(jorgensen_b$V2), ]

xb_jorgensen <- sort(jorgensen_b$V1)
yb_jorgensen <- jorgensen_b$V2
```

## Startfunktion

Um die bestmögliche Anpassung an den vorliegenden Datenpunkten darzustellen wurde mithilfe einer Funktion der beste Startparameter mit dem kleinsten Fehler ermittelt, da die Anpassung stark von den initialen Startwerten abhängig ist.

```
find_best_start_2parameter <- function(p, q, max_beta, max_eta, steps_beta,
                                         steps_eta, fitting_function) {
```

```

best_errors <- numeric() # Vektor für Fehlerwerte
best_starts <- matrix(nrow = 0, ncol = 2) # Matrix für Startparameter

for (beta in seq(0, max_beta, by = steps_beta)) {
  for (eta in seq(0, max_eta, by = steps_eta)) {
    start_params <- c(beta, eta)

    # Schätze die Parameter mit den aktuellen Startparametern
    if(identical(as.character(substitute(fitting_function)), "getstuexp2")){
      output <- capture.output({
        result <- getstuexp2(
          p = p, q = q, start = start_params,
          show.output = TRUE, plot = FALSE, wert1 = 2
        )
      })
    }
    else{
      output <- capture.output({
        result <- fitting_function(p = p, q = q, start = start_params,
          show.output = TRUE, plot = FALSE)
      })
    }

    # Berechne den Fehler (jorgensen_r_1$value) für die aktuellen Startparameter
    current_error <- as.numeric(gsub("\\[1\\]\\s+", "", output[5]))

    # Speichere Fehler und die Startparameter, wenn der Fehler nicht NA ist
    if (!is.na(current_error)) {
      best_errors <- c(best_errors, current_error)
      best_starts <- rbind(best_starts, start_params)
    }
  }
}

# Finde den Index des kleinsten Fehlers (ignoriere NA-Werte)
best_index <- which.min(best_errors)

# Wähle den besten Startparameter mit dem kleinsten Fehler aus
best_start <- best_starts[best_index, ]

# Gib den besten Startparameter und den entsprechenden Fehler aus
cat("Bester Startparameter:", best_start, "\n")
cat("Bester Fehler:", best_errors[best_index], "\n")

return(best_start)
}

find_best_start_3parameter <- function(p, q, max_shape1 = 10, max_shape2 = 10,
  max_scale = 10, steps_shape1, steps_shape2,
  steps_scale, fitting_function) {

```

```

best_errors <- numeric() # Vektor für Fehlerwerte
best_starts <- matrix(nrow = 0, ncol = 3) # Matrix für Startparameter

for (shape1 in seq(0, max_shape1, by = steps_shape1)) {
  for (shape2 in seq(0, max_shape2, by = steps_shape2)) {
    for (scale in seq(0, max_shape1, by = steps_scale)) {
      start_params <- c(shape1, shape2, scale)

      # Schätze die Parameter mit den aktuellen Startparametern
      if(identical(as.character(substitute(fitting_function)), "getstuexp3")){
        output <- capture.output({
          result <- getstuexp3(
            p = p, q = q, start = start_params,
            show.output = TRUE, plot = FALSE, wert1 = 2, wert2 = 6)
        })
      }
      else{
        output <- capture.output({
          result <- fitting_function(p = p, q = q, start = start_params,
            show.output = TRUE, plot = FALSE)
        })
      }
      # Berechne den Fehler (jorgensen_r_1$value) für die aktuellen Startparameter
      current_error <- as.numeric(gsub("\\[1\\]\\s+", "", output[5]))

      # Speichere Fehler und die Startparameter, wenn der Fehler nicht NA ist
      if (!is.na(current_error)) {
        best_errors <- c(best_errors, current_error)
        best_starts <- rbind(best_starts, start_params)
      }
    }
  }
}

# Finde den Index des kleinsten Fehlers (ignoriere NA-Werte)
best_index <- which.min(best_errors)

# Wähle den besten Startparameter mit dem kleinsten Fehler aus
best_start <- best_starts[best_index, ]

# Gib den besten Startparameter und den entsprechenden Fehler aus
cat("Bester Startparameter:", best_start, "\n")
cat("Bester Fehler:", best_errors[best_index], "\n")

return(best_start)
}

find_best_start_4parameter <- function(p, q, max_shape, max_scale, max_rate,
  max_mix, steps_shape, steps_scale,

```

```

                                steps_rate, steps_mix, fitting_function) {

best_errors <- numeric() # Vektor für Fehlerwerte
best_starts <- matrix(nrow = 0, ncol = 4) # Matrix für Startparameter

for (shape in seq(0, max_shape, by = steps_shape)) {
  for (scale in seq(0, max_scale, by = steps_scale)) {
    for (rate in seq(0, max_rate, by = steps_rate)) {
      for (mix in seq(0, max_mix, by = steps_mix)) {
        start_params <- c(shape, scale, rate, mix)

# Schätze die Weibull-Parameter mit den aktuellen Startparametern
        output <- capture.output({
          result <- fitting_function(p = p, q = q, start = start_params,
                                     show.output = TRUE, plot = FALSE)
        })

# Berechne den Fehler (jorgensen_r_1$value) für die aktuellen Startparameter
        current_error <- as.numeric(gsub("\\[1\\]\\s+", "", output[5]))

# Speichere Fehler und die Startparameter, wenn der Fehler nicht NA ist
        if (!is.na(current_error)) {
          best_errors <- c(best_errors, current_error)
          best_starts <- rbind(best_starts, start_params)
        }
      }
    }
  }
}

# Finde den Index des kleinsten Fehlers (ignoriere NA-Werte)
best_index <- which.min(best_errors)

# Wähle den besten Startparameter mit dem kleinsten Fehler aus
best_start <- best_starts[best_index, ]

# Gib den besten Startparameter und den entsprechenden Fehler aus
cat("Bester Startparameter:", best_start, "\n")
cat("Bester Fehler:", best_errors[best_index], "\n")

return(best_start)
}

```

## Datenanpassung an die Daten von Jorgensen

### Weibullverteilung

```

# Weibullverteilung
source("C:/Users/Chau/Documents/Masterarbeit/R Funktionen/getweibullpar.R")

best_jorgensen_r_1 <- find_best_start_2parameter(p = yr_jorgensen/100, q = xr_jorgensen,
                                                max_beta = 10, max_eta = 10,

```

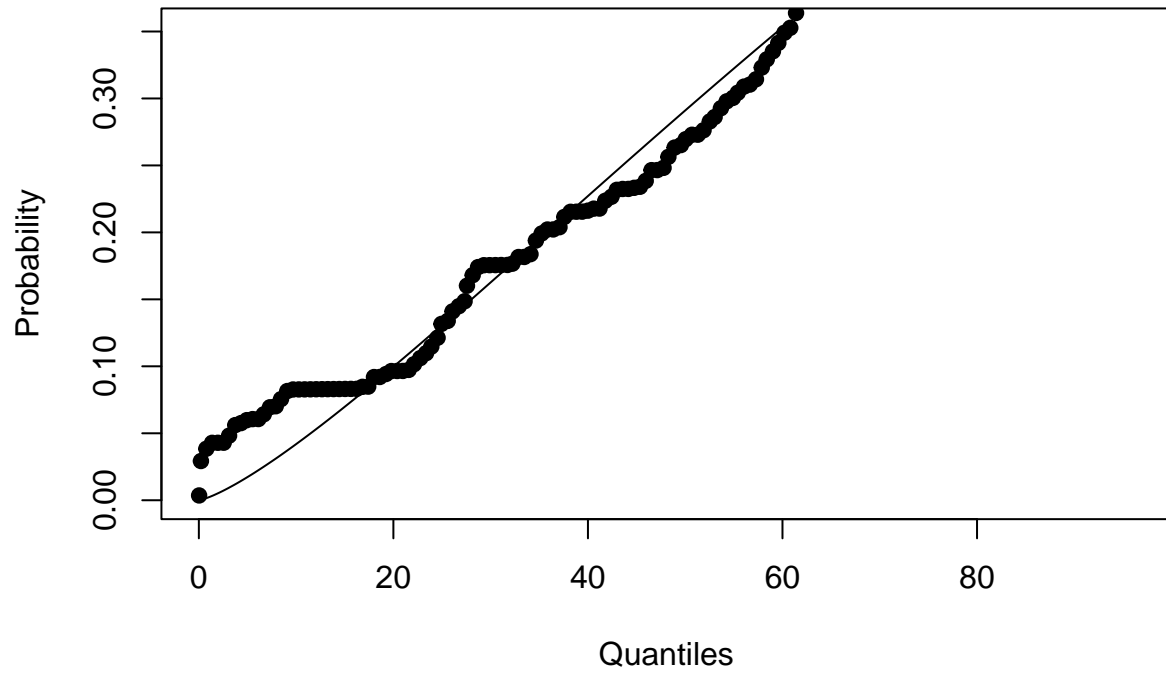
```
steps_beta = 1, steps_eta = 1,  
fitting_function = getweibullpar)
```

```
## Bester Startparameter: 2 5  
## Bester Fehler: 2.18233e-06
```

```
jorgensen_r_1 <- getweibullpar(  
  p = yr_jorgensen/100,  
  q = xr_jorgensen,  
  start = best_jorgensen_r_1,  
  show.output = TRUE,  
  plot = TRUE  
)
```

```
## $par  
## [1] 1.294875 114.000721  
##  
## $value  
## [1] 2.18233e-06  
##  
## $counts  
## function gradient  
## 77 77  
##  
## $convergence  
## [1] 0  
##  
## $message  
## [1] "CONVERGENCE: REL_REDUCTION_OF_F <= FACTR*EPSMCH"
```

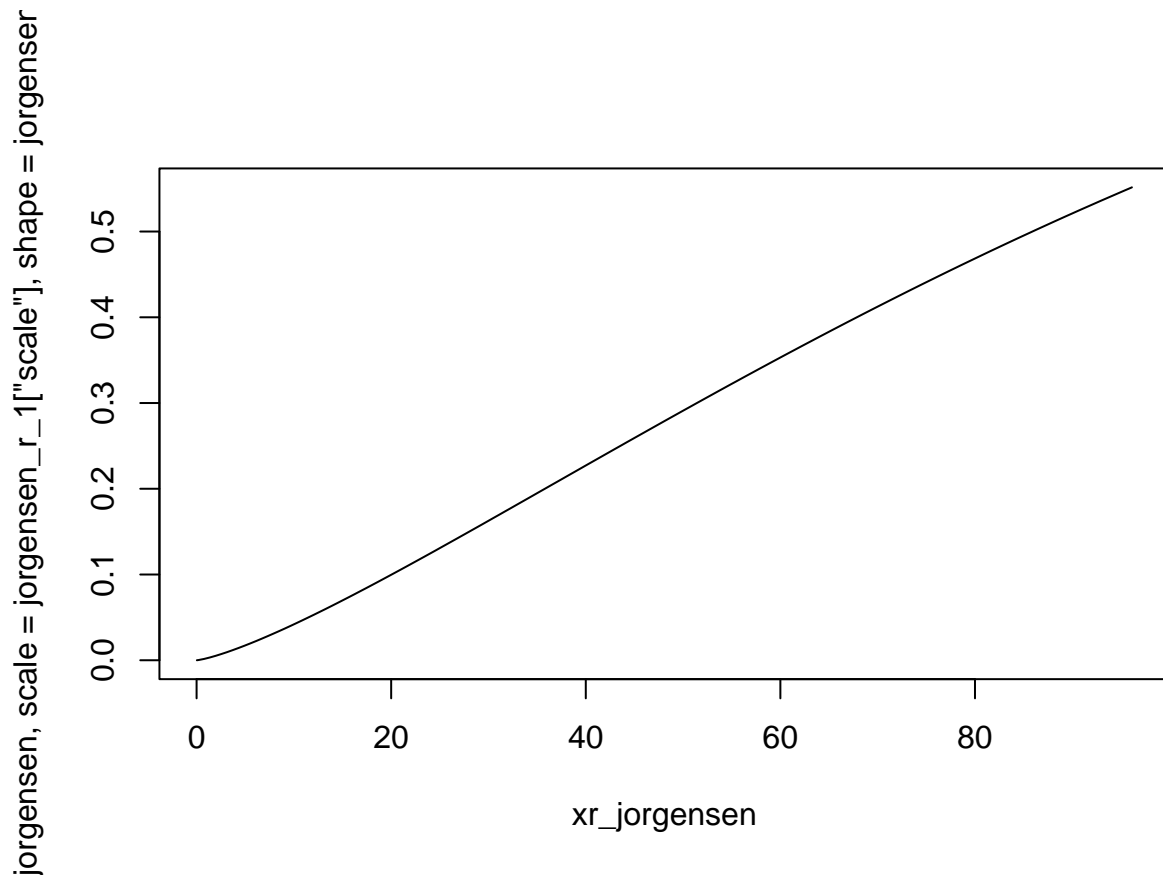
**Weibull (shape = 1.29, scale = 114)**



```
jorgensen_r_1
```

```
##      shape      scale
##  1.294875 114.000721
```

```
plot(xr_jorgensen,
     pweibull(xr_jorgensen,
              scale = jorgensen_r_1["scale"],
              shape = jorgensen_r_1["shape"]), type = "l")
```



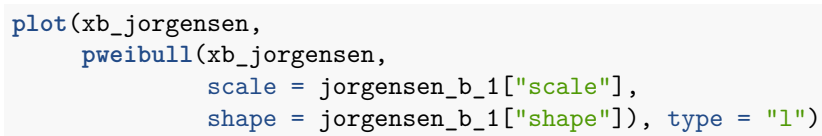
```
best_jorgensen_b_1 <- find_best_start_2parameter(p = yb_jorgensen/100, q = xb_jorgensen,
max_beta = 10, max_eta = 10,
steps_beta = 1, steps_eta = 1,
fitting_function = getweibullpar)
```

```
## Bester Startparameter: 2 2
## Bester Fehler: 3.334577e-06
```

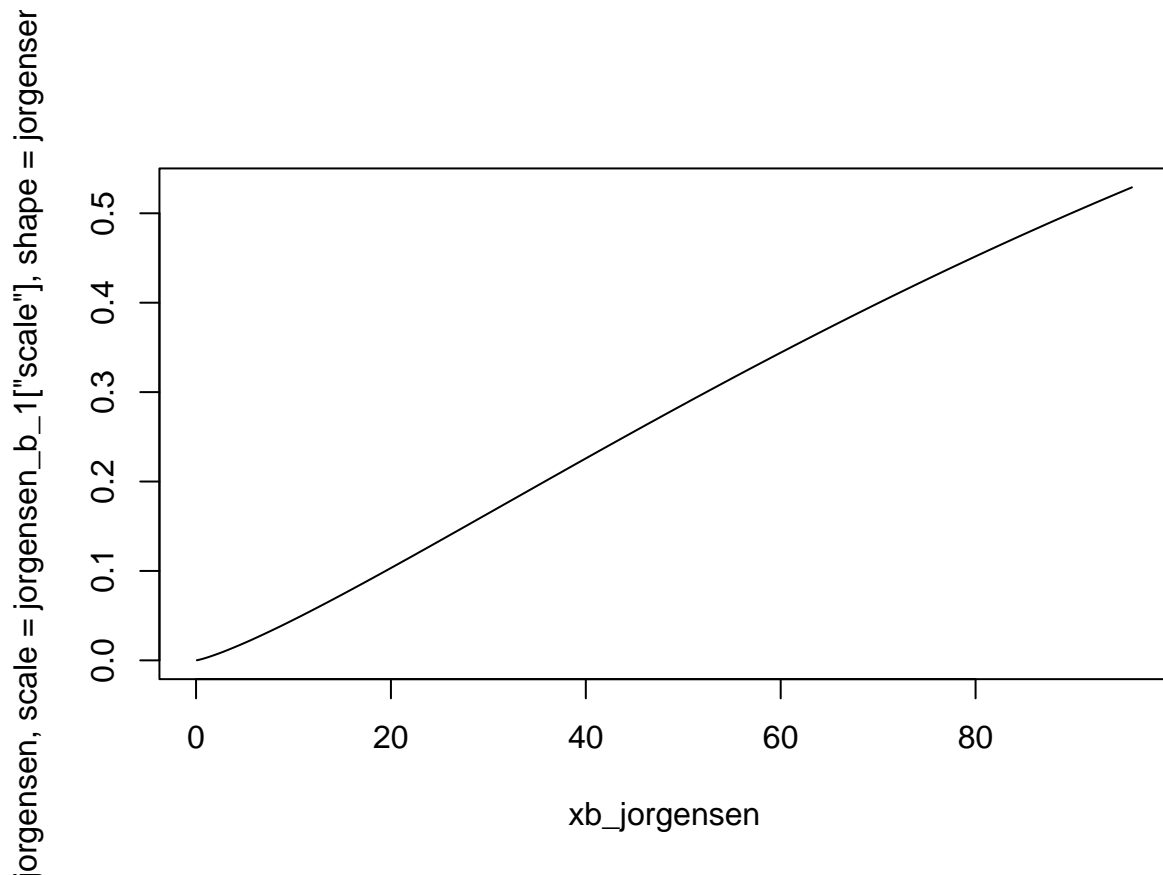
```
jorgensen_b_1 <- getweibullpar(
  p = yb_jorgensen/100,
  q = xb_jorgensen,
  start = best_jorgensen_b_1,
  show.output = TRUE,
  plot = TRUE
)
```

```
## $par
## [1] 1.231547 120.941739
##
## $value
## [1] 3.334577e-06
##
## $counts
## function gradient
##      57      57
##
## $convergence
```

```
plot(xb_jorgensen,
     pweibull(xb_jorgensen,
              scale = jorgensen_b_1["scale"],
              shape = jorgensen_b_1["shape"]), type = "l")
```







## Exponentiierte Weibullverteilung

```
# exponentiierte Weibullverteilung
source("C:/Users/Chau/Documents/Masterarbeit/R Funktionen/getweibpar.R")

best_weibbull_xr_jorgensen <- find_best_start_2parameter(p = yr_jorgensen/100,
                                                         q = xr_jorgensen,
                                                         max_beta = 10,
                                                         max_eta = 10,
                                                         steps_beta = 1,
                                                         steps_eta = 1,
                                                         fitting_function = getweibpar)

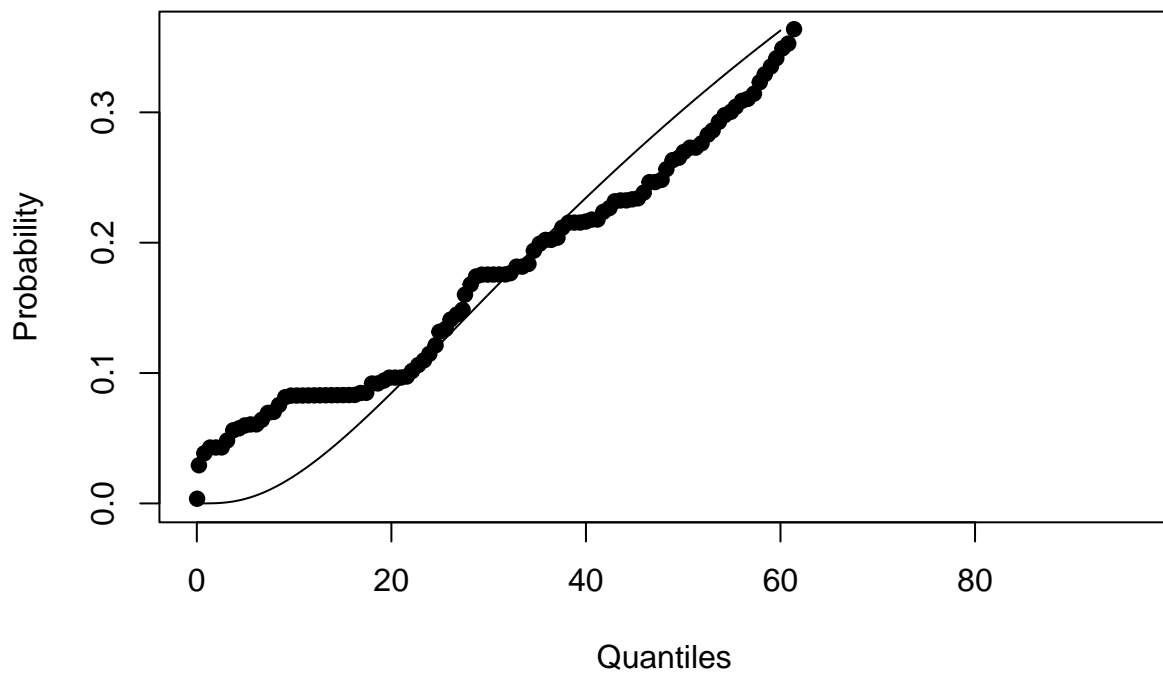
## Bester Startparameter: 5 0
## Bester Fehler: 4.358661e-06

weibbull_xr_jorgensen <- getweibpar(
  p = yr_jorgensen/100,
  q = xr_jorgensen,
  start = best_weibbull_xr_jorgensen,
  show.output = TRUE,
  plot = TRUE
)

## $par
## [1] 0.2845173 24.5051353
```

```
##  
## $value  
## [1] 4.358661e-06  
##  
## $counts  
## function gradient  
##      38      38  
##  
## $convergence  
## [1] 0  
##  
## $message  
## [1] "CONVERGENCE: REL_REDUCTION_OF_F <= FACTR*EPSMCH"
```

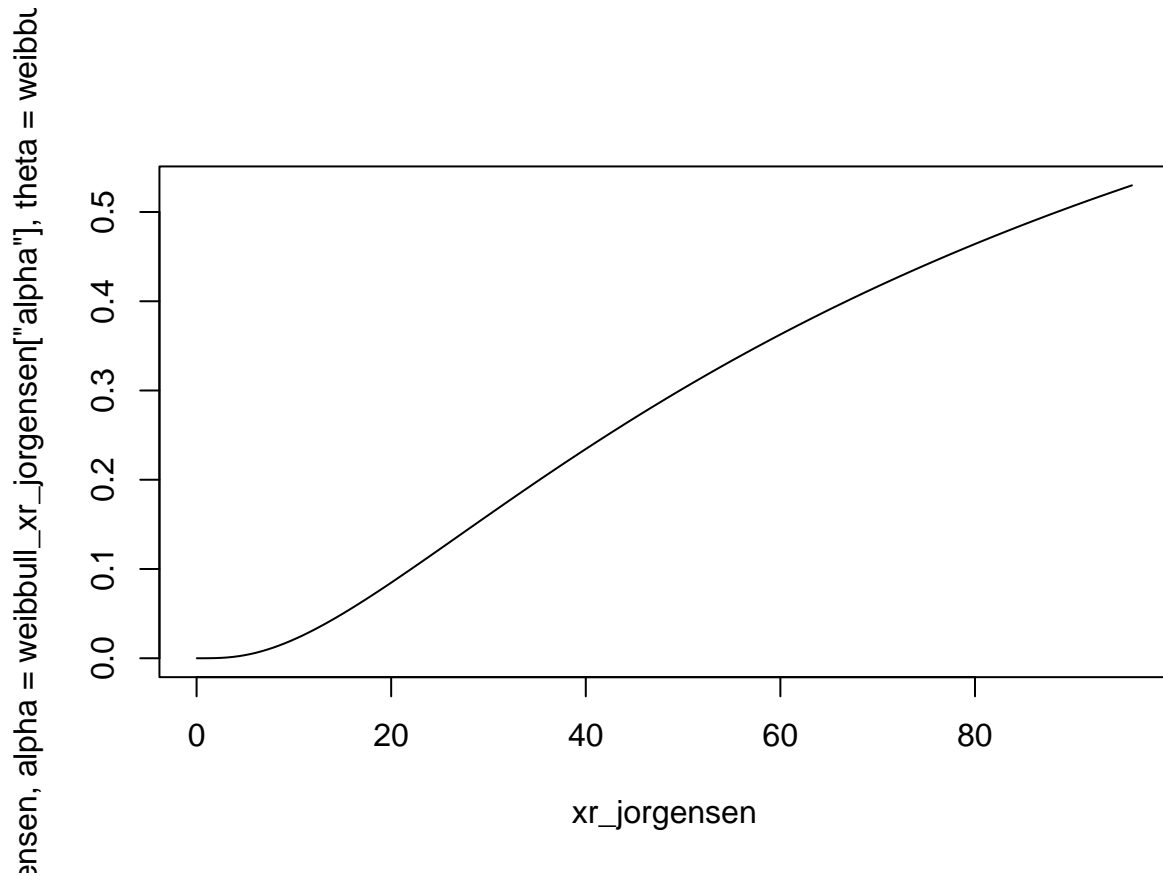
**exp. Weibull (alpha = 0.285, theta = 24.5)**

[illegible]

weibbull\_xr\_jorgensen

```
##      alpha      theta
## 0.2845173 24.5051353
```

```
plot(xr_jorgensen,
     pexpo.weibull(xr_jorgensen,
                   alpha = weibull_xr_jorgensen ["alpha"],
                   theta = weibull_xr_jorgensen ["theta"]), type = "l")
```



```
best_weibull_xb_jorgensen <- find_best_start_2parameter(p = yb_jorgensen/100,
                                                       q = xb_jorgensen,
                                                       max_beta = 10,
                                                       max_eta = 10,
                                                       steps_beta = 1,
                                                       steps_eta = 1,
                                                       fitting_function = getweibpar)
```

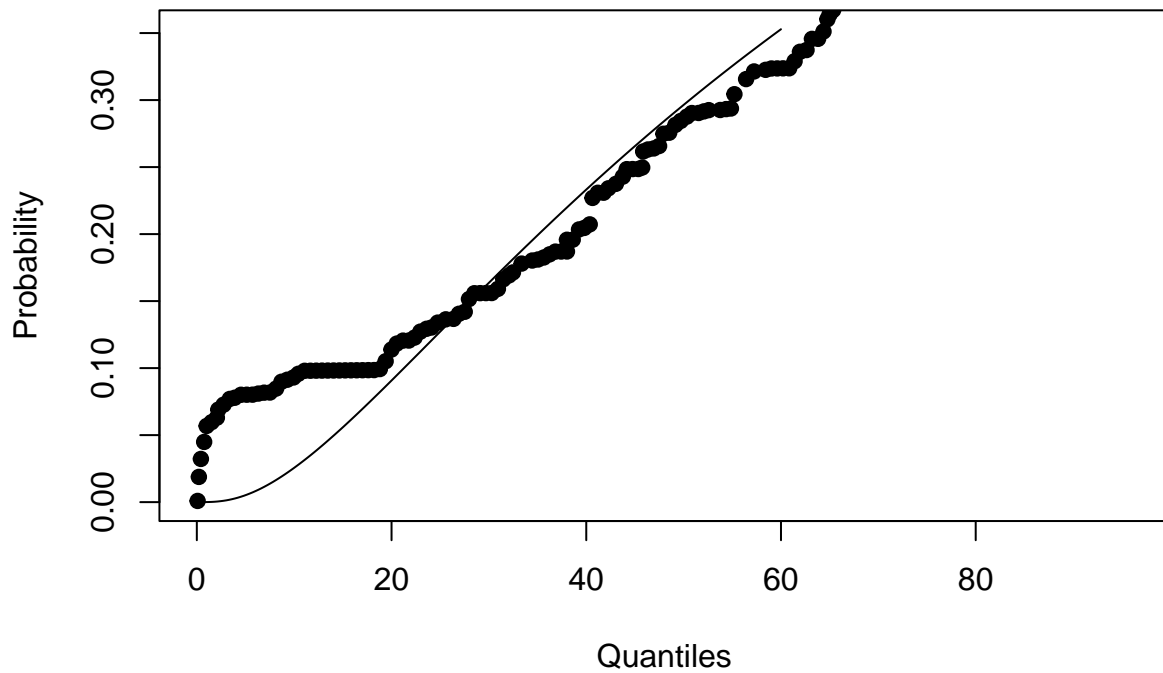
```
## Bester Startparameter: 0 4
## Bester Fehler: 6.085128e-06
```

```
weibull_xb_jorgensen <- getweibpar(
  p = yb_jorgensen/100,
  q = xb_jorgensen,
  start = best_weibull_xb_jorgensen,
  show.output = TRUE,
  plot = TRUE
)
```

```
## $par
## [1] 0.2750718 22.2315848
##
## $value
## [1] 6.085128e-06
##
## $counts
## function gradient
```

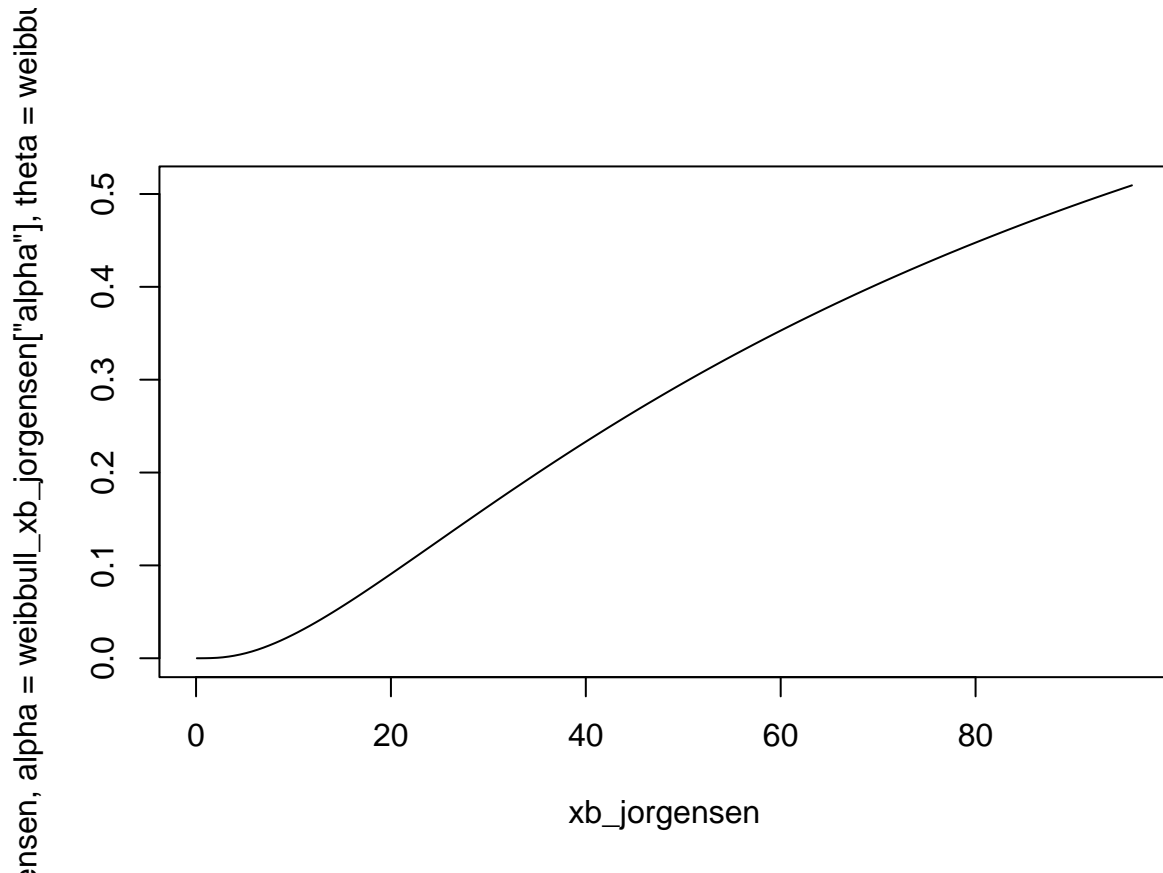
```
##          28          28
##
## $convergence
## [1] 0
##
## $message
## [1] "CONVERGENCE: REL_REDUCTION_OF_F <= FACTR*EPSMCH"
```

**exp. Weibull (alpha = 0.275, theta = 22.2)**

[illegible]

```
##      alpha      theta
## 0.2750718 22.2315848

plot(xb_jorgensen,
     pexpo.weibull(xb_jorgensen,
                   alpha = weibull_xb_jorgensen ["alpha"],
                   theta = weibull_xb_jorgensen ["theta"]), type = "l")
```



## Mischung aus Weibull- und Exponentialverteilung

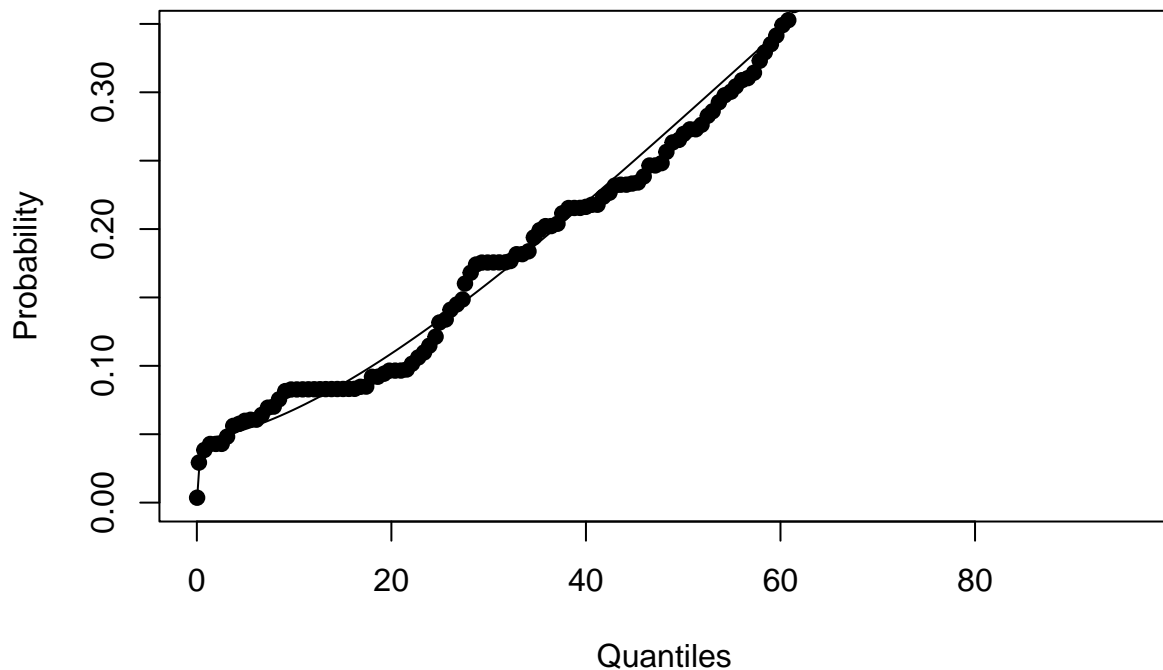
```
# Mischung aus Weibull- und Exponentialverteilung
source("C:/Users/Chau/Documents/Masterarbeit/R Funktionen/getweibex.R")

best_weibex_xr_jorgensen <- find_best_start_4parameter(p = yr_jorgensen/100,
  q = xr_jorgensen,
  max_shape = 1,
  max_scale = 100,
  max_rate = 0.7,
  max_mix = 1,
  steps_shape = 0.1,
  steps_scale = 20,
  steps_rate = 0.1,
  steps_mix = 0.1,
  fitting_function = getweibex)

## Bester Startparameter: 0.5 100 0.6 0.2
## Bester Fehler: 9.427333e-07

weibex_xr_jorgensen <- getweibex(
  p = yr_jorgensen/100,
  q = xr_jorgensen,
  start = best_weibex_xr_jorgensen, # c(0.5, 60, 0.4, 0.1),
  show.output = TRUE,
  plot = TRUE
```

**Weibull & Exponential (shape = 1.56, scale = 112, rate = 3.78, mix = 0.9)**

[illegible]

weibex\_xr\_jorgensen

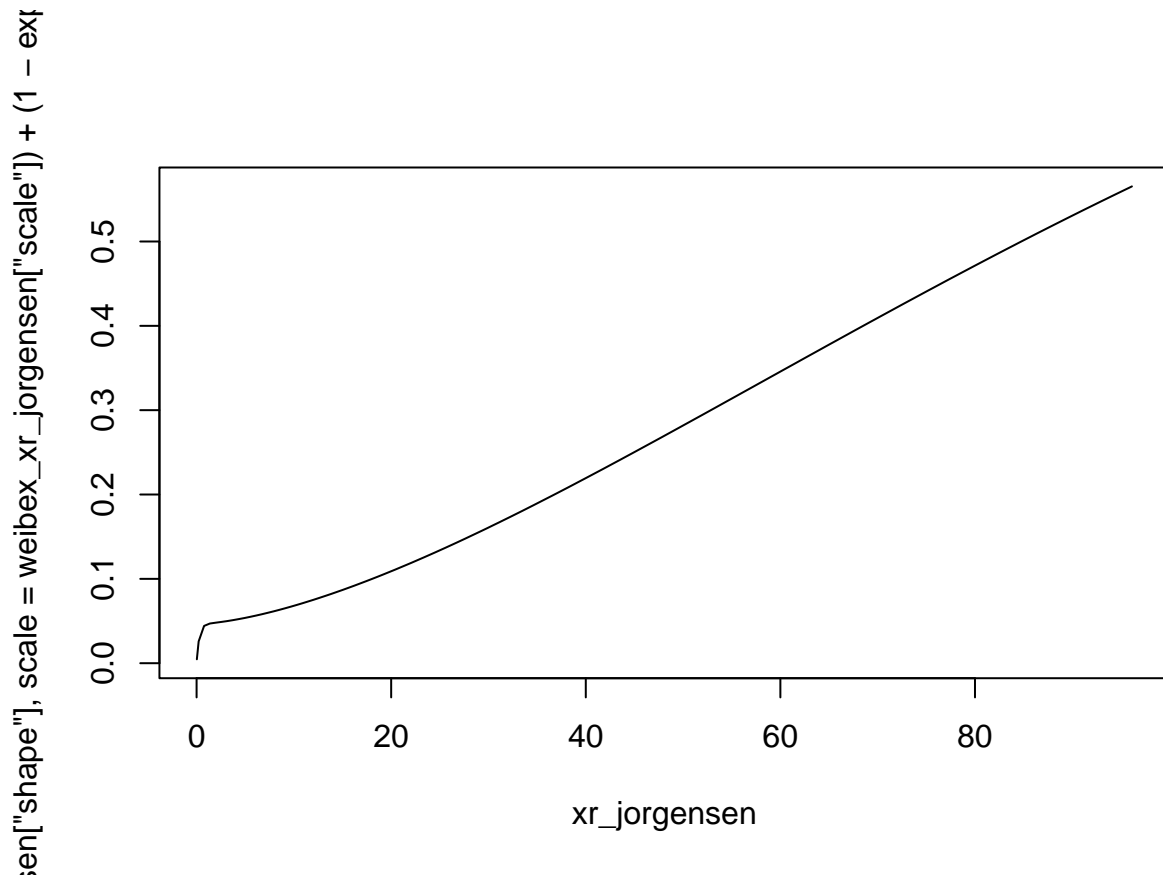
```
##      shape      scale      rate      mix
##  1.558657 112.219015   3.780190   3.025024

# Wenn Ergebnisse aus weibex_xr1_1 von Funktion abgelesen
plot(xr_jorgensen,
      (exp(weibex_xr_jorgensen["mix"]) / ( 1 + exp(weibex_xr_jorgensen["mix"]))) *
      stats::pweibull(q = xr_jorgensen,
```

```

        shape = weibex_xr_jorgensen["shape"],
        scale = weibex_xr_jorgensen["scale"]) +
(1 - exp(weibex_xr_jorgensen["mix"])) / (1 + exp(weibex_xr_jorgensen["mix"])) *
stats::pexp(q = xr_jorgensen,
            rate = weibex_xr_jorgensen["rate"])),
type = "l")

```



```

best_weibex_xb_jorgensen <- find_best_start_4parameter(p = yb_jorgensen/100,
                                                    q = xb_jorgensen,
                                                    max_shape = 1,
                                                    max_scale = 100,
                                                    max_rate = 0.7,
                                                    max_mix = 1,
                                                    steps_shape = 0.1,
                                                    steps_scale = 20,
                                                    steps_rate = 0.1,
                                                    steps_mix = 0.1,
                                                    fitting_function = getweibex)

```

```

## Bester Startparameter: 0.8 100 0.7 0.8
## Bester Fehler: 5.537916e-07

```

```

weibex_xb_jorgensen <- getweibex(
  p = yb_jorgensen/100,
  q = xb_jorgensen,
  start = best_weibex_xb_jorgensen, # c(0.5, 60, 0.7, 0.1),
  show.output = TRUE,

```

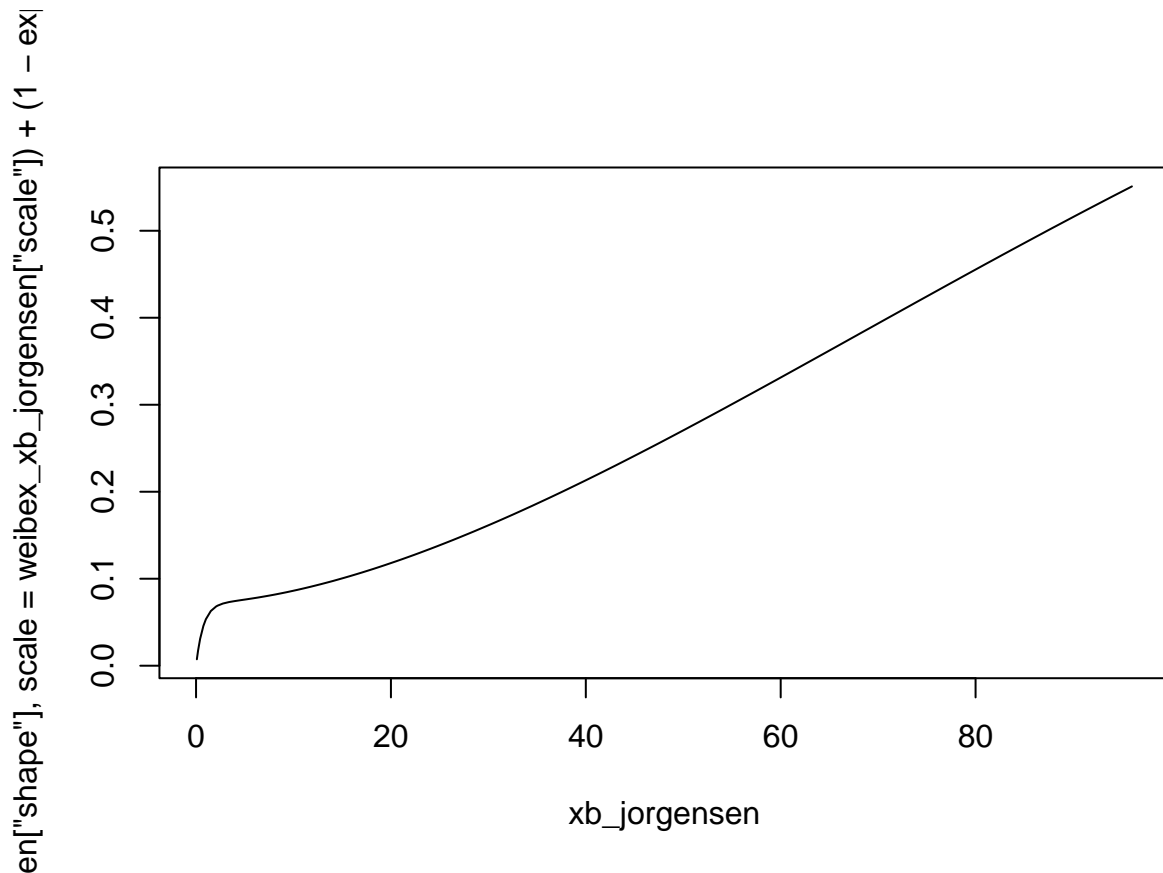




```

        shape = weibex_xb_jorgensen["shape"],
        scale = weibex_xb_jorgensen["scale"]) +
(1 - exp(weibex_xb_jorgensen["mix"])) / (1 + exp(weibex_xb_jorgensen["mix"])) *
stats::pexp(q = xb_jorgensen,
            rate = weibex_xb_jorgensen["rate"])),
type = "l")

```



## Mischung von exponentiierter Weibull- und Exponentialverteilung

```

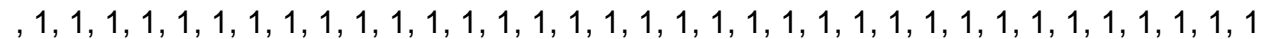
# Mischung von exponentiierter Weibull- und Exponentialverteilung
source("C:/Users/Chau/Documents/Masterarbeit/R Funktionen/get2weibex.R")

best_weibex2_xr_jorgensen <- find_best_start_4parameter(p = yr_jorgensen/100,
    q = xr_jorgensen,
    max_shape = 1,
    max_scale = 100,
    max_rate = 0.7,
    max_mix = 1,
    steps_shape = 0.1,
    steps_scale = 20,
    steps_rate = 0.1,
    steps_mix = 0.1,
    fitting_function = get2weibex)

## Bester Startparameter: 0.1 0 0.4 0.4

```

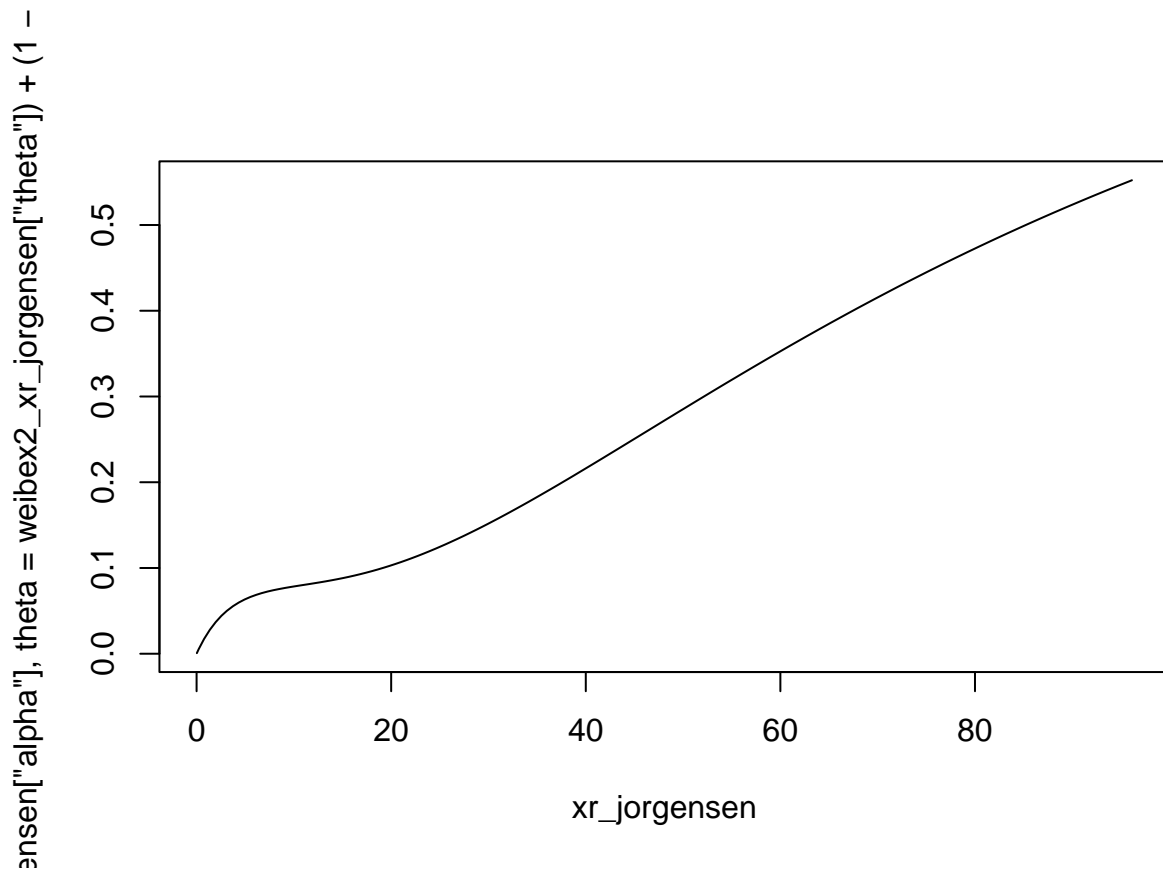
```
weibex2_xr_jorgensen <- get2weibex(  
  p = yr_jorgensen/100,  
  q = xr_jorgensen,  
  start = best_weibex2_xr_jorgensen,  
  show.output = TRUE,  
  plot = TRUE  
)
```



```
weibex2_xr_jorgensen
```

```
##      alpha      theta      rate      mix
## 0.3180856 47.5324993 0.3130429 2.4378076
```

```
plot(xr_jorgensen,
      (exp(weibex2_xr_jorgensen["mix"]) / (1 + exp(weibex2_xr_jorgensen["mix"]))) *
        reliaR::pexpo.weibull(q = xr_jorgensen,
                              alpha = weibex2_xr_jorgensen["alpha"],
                              theta = weibex2_xr_jorgensen["theta"]) +
      (1 - exp(weibex2_xr_jorgensen["mix"]) / (1 + exp(weibex2_xr_jorgensen["mix"]))) *
        stats::pexp(q = xr_jorgensen,
                    rate = weibex2_xr_jorgensen["rate"])),
      type = "l")
```



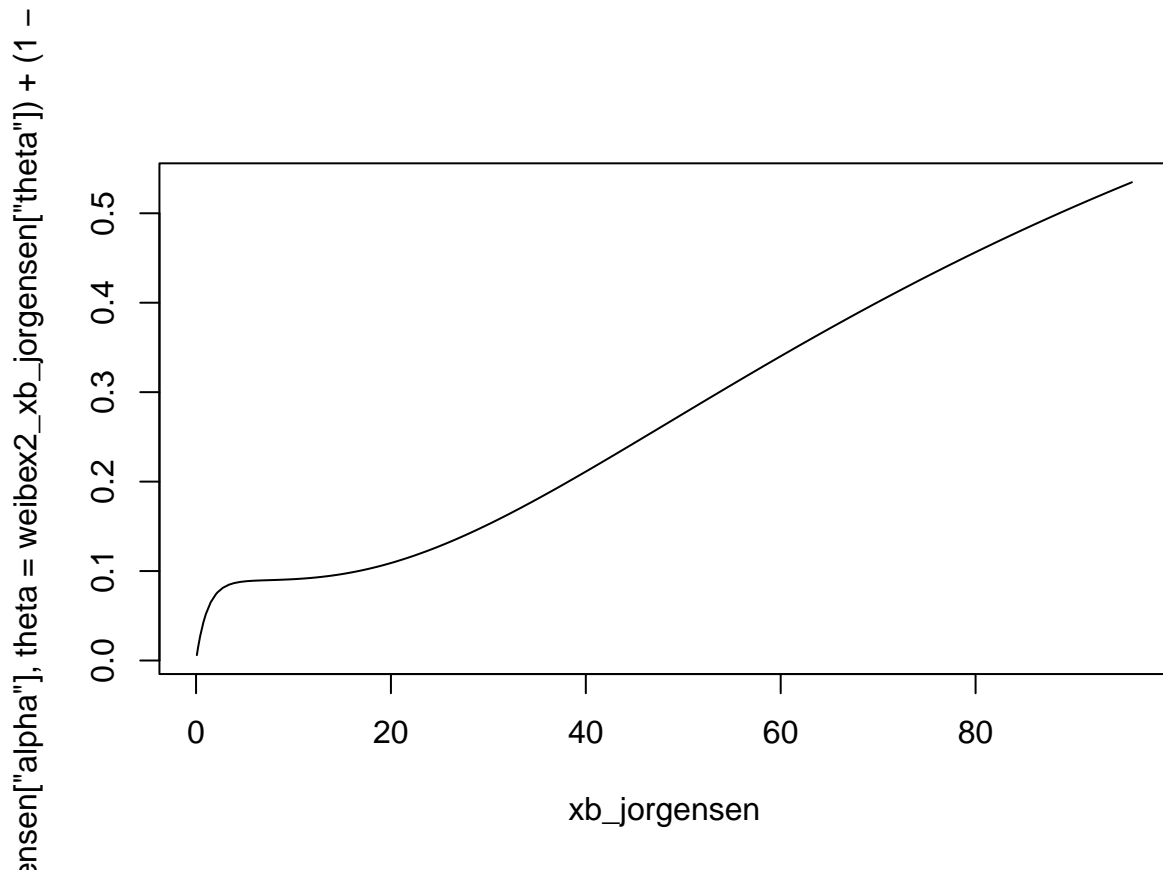
```
best_weibex2_xb_jorgensen <- find_best_start_4parameter(p = yb_jorgensen/100,
                                                         q = xb_jorgensen,
                                                         max_shape = 1,
                                                         max_scale = 100,
                                                         max_rate = 0.7,
                                                         max_mix = 1,
                                                         steps_shape = 0.1,
                                                         steps_scale = 20,
                                                         steps_rate = 0.1,
                                                         steps_mix = 0.1,
                                                         fitting_function = get2weibex)
```



```
weibex2_xb_jorgensen
```

```
##      alpha      theta      rate      mix
## 0.3157402 48.6334591 0.8606514 2.3178641
```

```
plot(xb_jorgensen,
     (exp(weibex2_xb_jorgensen["mix"]) / (1 + exp(weibex2_xb_jorgensen["mix"]))) *
       reliaR::pexpo.weibull(q = xb_jorgensen,
                             alpha = weibex2_xb_jorgensen["alpha"],
                             theta = weibex2_xb_jorgensen["theta"]) +
     (1 - exp(weibex2_xb_jorgensen["mix"]) / (1 + exp(weibex2_xb_jorgensen["mix"]))) *
       stats::pexp(q = xb_jorgensen,
                   rate = weibex2_xb_jorgensen["rate"])),
     type = "l")
```



### Exponentiierte Weibullverteilung ohne Lambda = 1

```
# exponentiierte Weibullverteilung ohne Lambda = 1
source("C:/Users/Chau/Documents/Masterarbeit/R Funktionen/getexpweib.R")

best_expweib_xr_jorgensen <- find_best_start_3parameter(p = yr_jorgensen/100,
                                                         q = xr_jorgensen,
                                                         max_shape1 = 10,
                                                         max_shape2 = 10,
                                                         max_scale = 10,
                                                         steps_shape1 = 1,
```

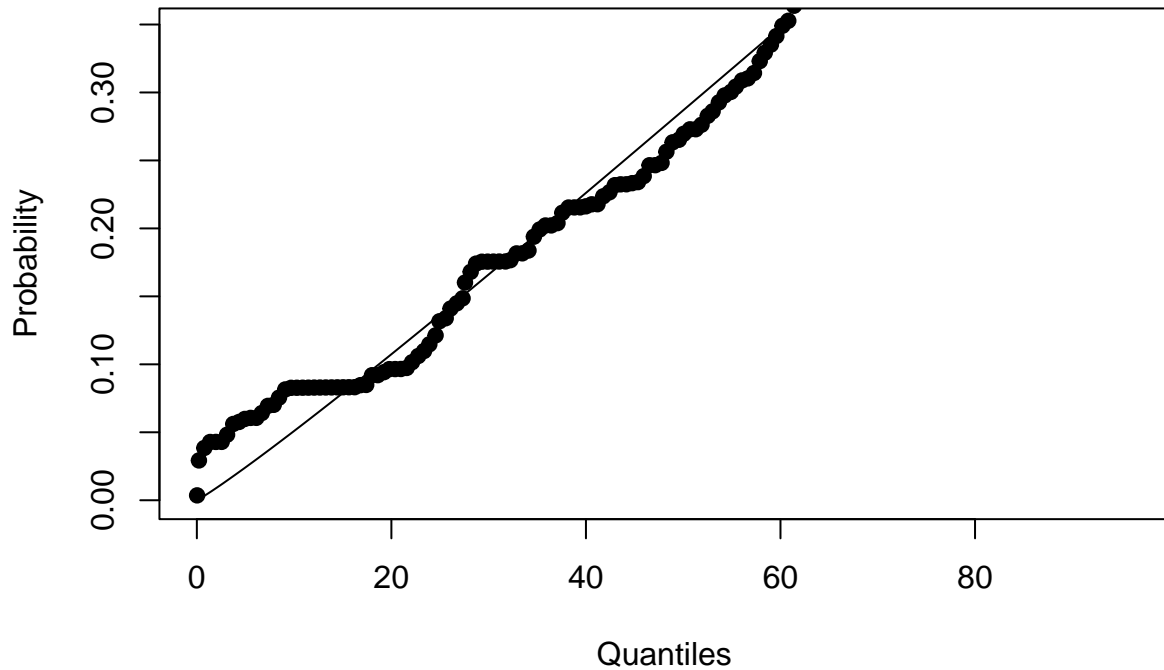
```
steps_shape2 = 1,  
steps_scale = 1,  
fitting_function = getexpweib)
```

```
## Bester Startparameter: 9 10 7  
## Bester Fehler: 1.799463e-06
```

```
expweib_xr_jorgensen <- getexpweib(  
  p = yr_jorgensen/100,  
  q = xr_jorgensen,  
  start = best_expweib_xr_jorgensen,  
  show.output = TRUE,  
  plot = TRUE  
)
```

```
## $par  
## [1] 158.9017624 3.1898719 0.3374211  
##  
## $value  
## [1] 1.799463e-06  
##  
## $counts  
## function gradient  
## 46 46  
##  
## $convergence  
## [1] 0  
##  
## $message  
## [1] "CONVERGENCE: REL_REDUCTION_OF_F <= FACTR*EPSMCH"
```

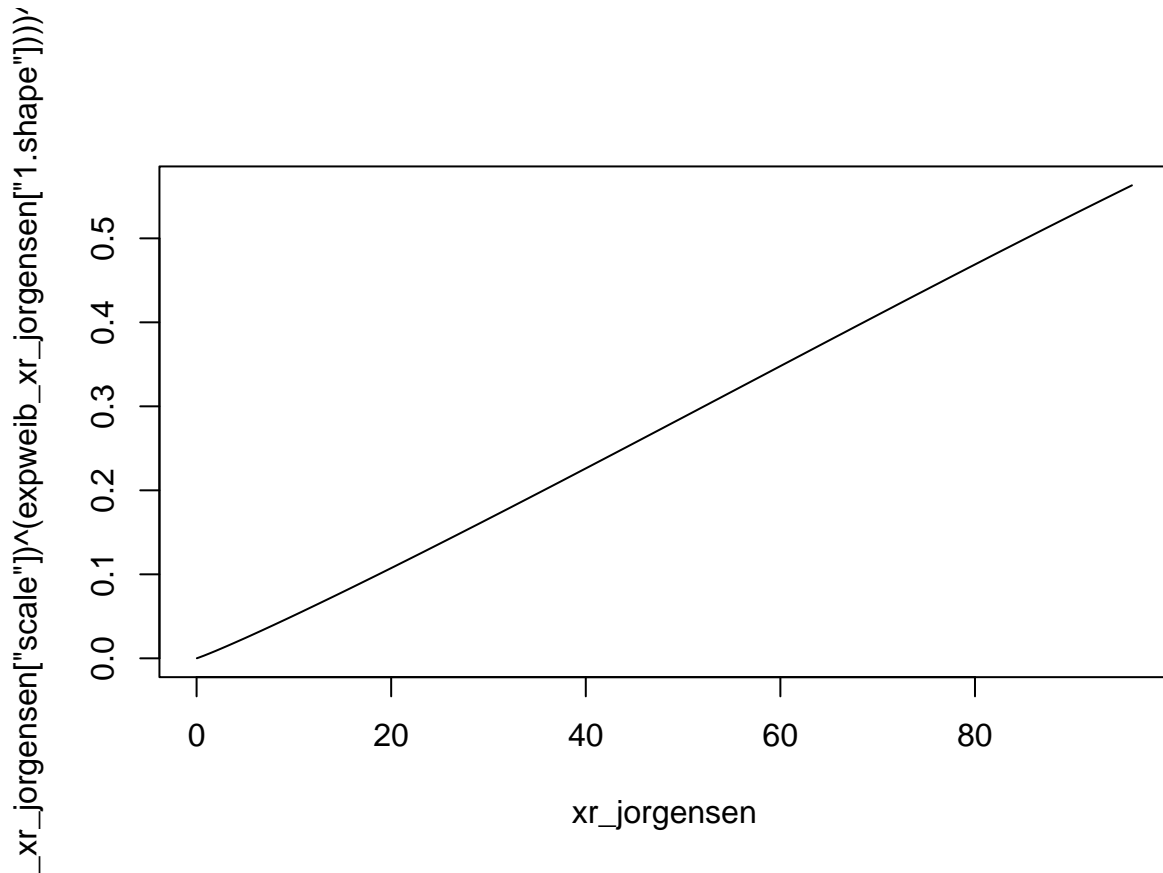
**exp. Weibull (scale = 159, 1.shape = 3.19, 2.shape = 0.337)**

[illegible]

expweib\_xr\_jorgensen

```
##          scale      1.shape      2.shape
## 158.9017624    3.1898719    0.3374211
```

```
plot(xr_jorgensen,
      (1 - exp(-(xr_jorgensen / expweib_xr_jorgensen["scale"]))^
        (expweib_xr_jorgensen["1.shape"]))) ^ (expweib_xr_jorgensen["2.shape"]),
      type = "l")
```



```
best_expweib_xb_jorgensen <- find_best_start_3parameter(p = yb_jorgensen/100,
                                                       q = xb_jorgensen,
                                                       max_shape1 = 10,
                                                       max_shape2 = 10,
                                                       max_scale = 10,
                                                       steps_shape1 = 1,
                                                       steps_shape2 = 1,
                                                       steps_scale = 1,
                                                       fitting_function = getexpweib)
```

```
## Bester Startparameter: 5 8 3
## Bester Fehler: 2.349781e-06
```

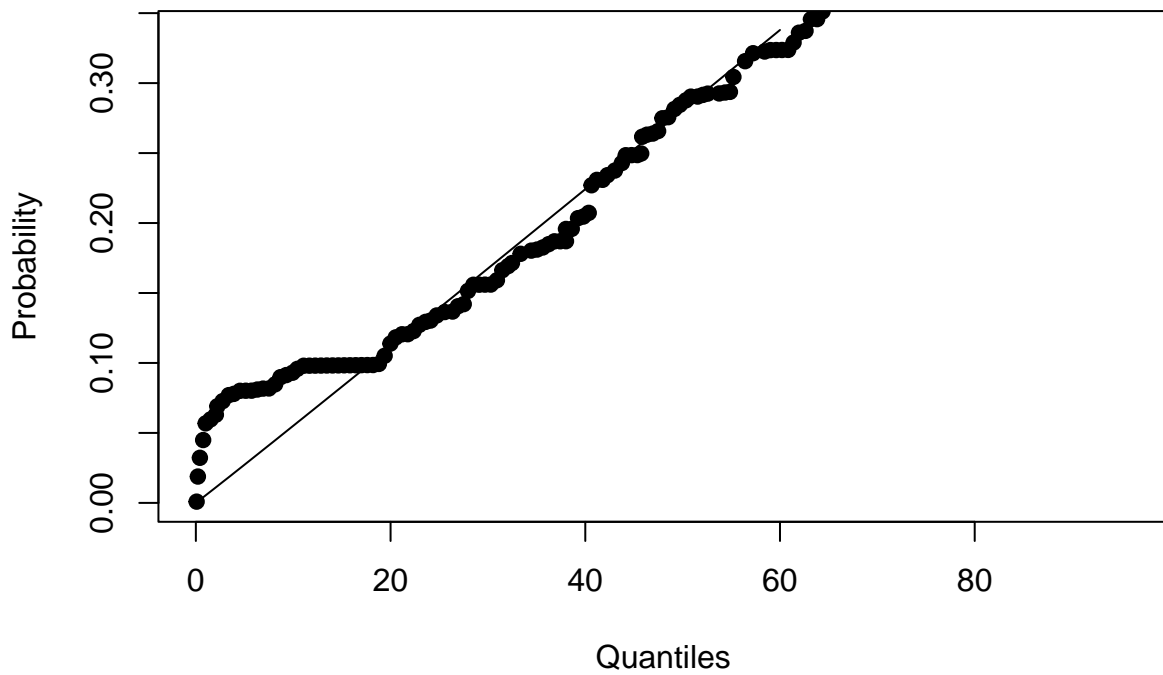
```
expweib_xb_jorgensen <- getexpweib(
  p = yb_jorgensen/100,
  q = xb_jorgensen,
  start = best_expweib_xb_jorgensen,
  show.output = TRUE,
  plot = TRUE
)
```

```
## $par
## [1] 175.1217978 6.1644571 0.1642775
##
## $value
## [1] 2.349781e-06
##
```



```
## $counts
## function gradient
##      57      57
##
## $convergence
## [1] 0
##
## $message
## [1] "CONVERGENCE: REL_REDUCTION_OF_F <= FACTR*EPSMCH"
```

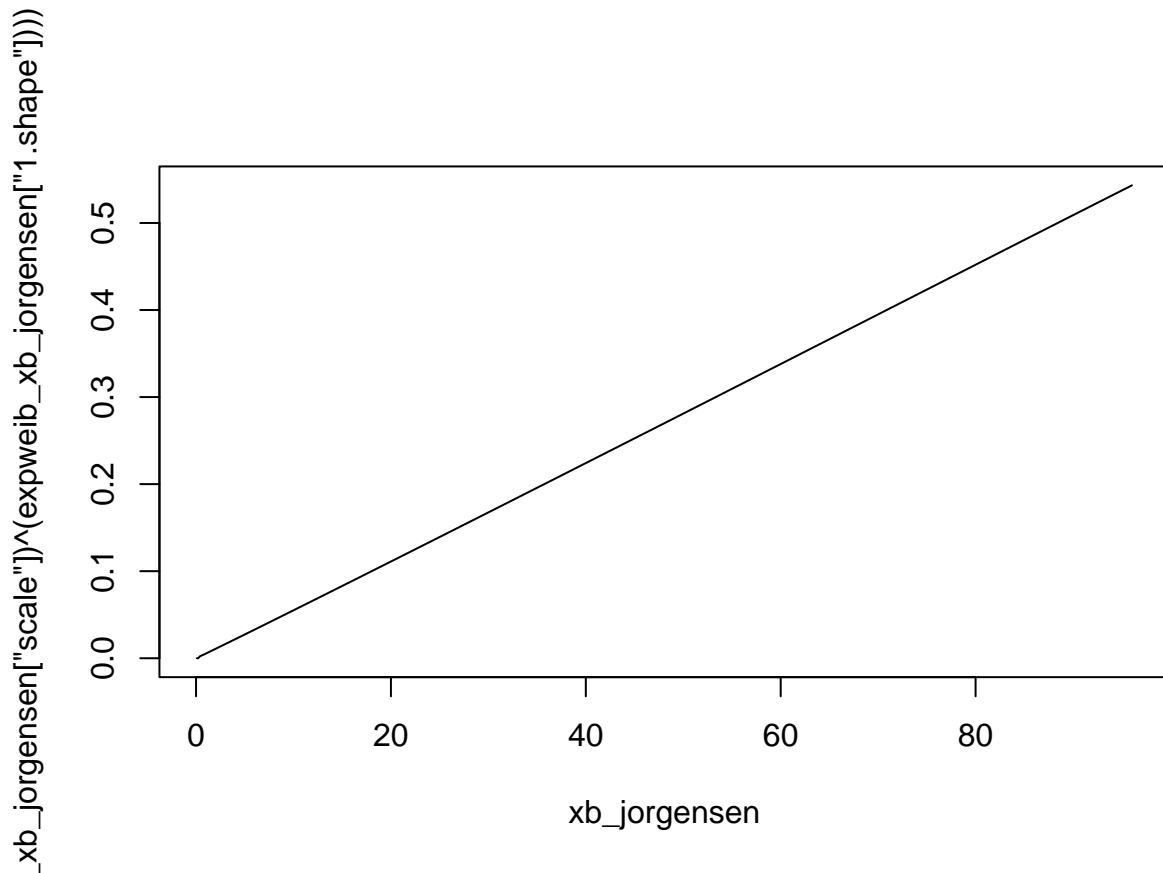
**exp. Weibull (scale = 175, 1.shape = 6.16, 2.shape = 0.164)**

[illegible]

expweib\_xb\_jorgensen

```
##          scale      1.shape      2.shape
## 175.1217978    6.1644571    0.1642775
```

```
plot(xb_jorgensen,
      (1 - exp(-(xb_jorgensen / expweib_xb_jorgensen["scale"]))^(expweib_xb_jorgensen["1.shape"])))^(expweib_xb_jorgensen["2.shape"])
      type = "l")
```



### 3-Stufige Exponentialverteilung

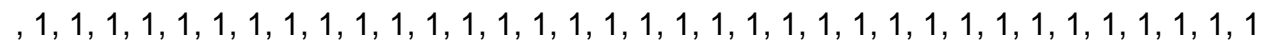
```
# 3-Stufige Exponentialverteilung
source("C:/Users/Chau/Documents/Masterarbeit/R Funktionen/getstuexp3.R")

best_stuexp3_xr_jorgensen <- find_best_start_3parameter(p = yr_jorgensen/100,
                                                       q = xr_jorgensen,
                                                       max_shape1 = 0.1,
                                                       max_shape2 = 0.1,
                                                       max_scale = 0.1,
                                                       steps_shape1 = 0.01,
                                                       steps_shape2 = 0.01,
                                                       steps_scale = 0.01,
                                                       fitting_function = getstuexp3)

## Bester Startparameter: 0.01 0.02 0.02
## Bester Fehler: 3.111139e-06

stuexp3_xr_jorgensen <- getstuexp3(
  p = yr_jorgensen/100,
  q = xr_jorgensen,
  start = best_stuexp3_xr_jorgensen,
  show.output = TRUE,
  plot = TRUE,
  wert1 = 2,
  wert2 = 6
)
```

3 stueckw. Exponential (1.para = 0.001, 2.para = 0.00331, 3.para = 0.0001)



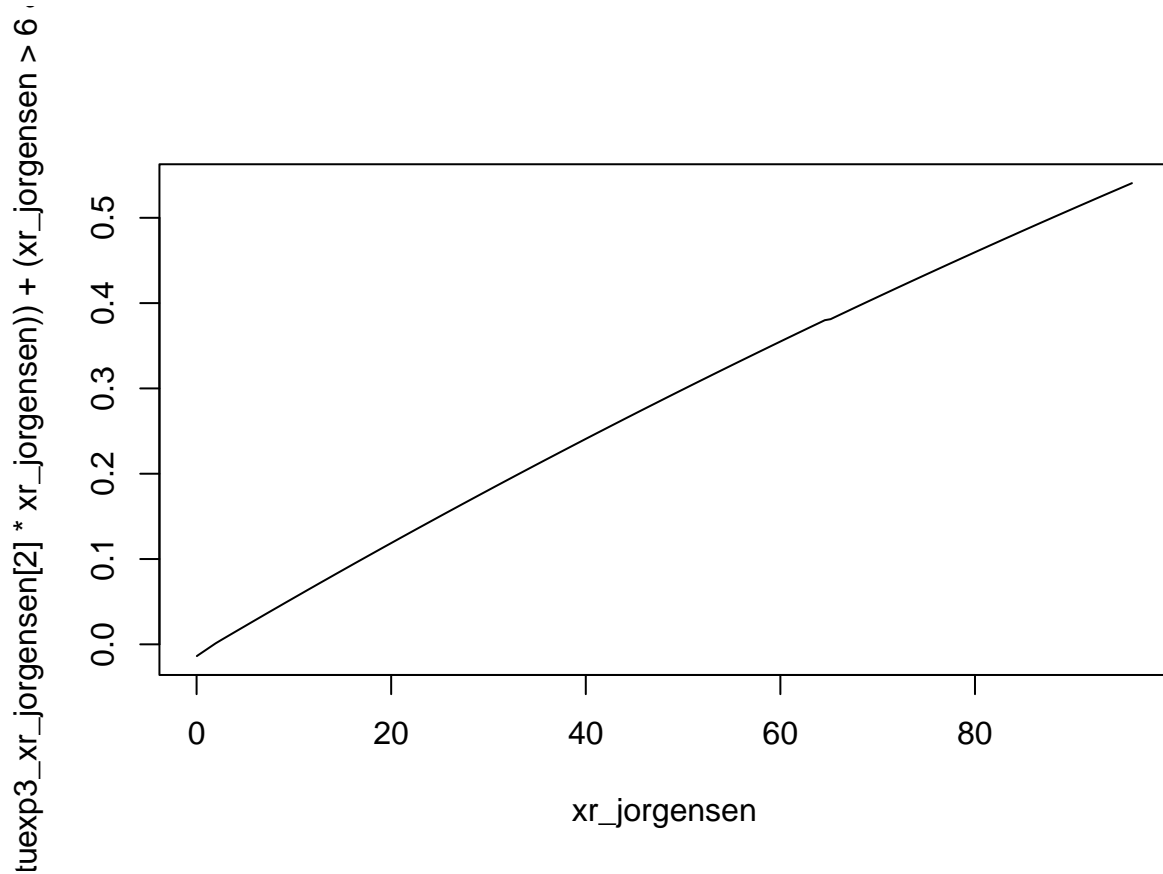
```
##          1.para      2.para      3.para
## 0.001000000 0.003313931 0.003442640

plot(xr_jorgensen,
      ((xr_jorgensen > 0 & xr_jorgensen <= 2) * (1 - exp(-stuexp3_xr_jorgensen[1] * xr_jorgensen)) +
        (xr_jorgensen > 2 & xr_jorgensen <= 6) *
        (1 - exp(-stuexp3_xr_jorgensen[1] * 2))) +
```

```

        (exp(-2 * stuexp3_xr_jorgensen[2]) -
          exp(-stuexp3_xr_jorgensen[2] * xr_jorgensen)) +
        (xr_jorgensen > 6 & xr_jorgensen <= 65 ) *
        (1 - exp(-stuexp3_xr_jorgensen[1] * 2)) +
        (exp(-2 * stuexp3_xr_jorgensen[2]) - exp(-6 * stuexp3_xr_jorgensen[2]
        (exp(-6 * stuexp3_xr_jorgensen[3]) -
          exp(-stuexp3_xr_jorgensen[3] * xr_jorgensen))),
type = "l")

```



```

best_stuexp3_xb_jorgensen <- find_best_start_3parameter(p = yb_jorgensen/100,
  q = xb_jorgensen,
  max_shape1 = 0.1,
  max_shape2 = 0.1,
  max_scale = 0.1,
  steps_shape1 = 0.01,
  steps_shape2 = 0.01,
  steps_scale = 0.01,
  fitting_function = getstuexp3)

```

```

## Bester Startparameter: 0 0.01 0.01
## Bester Fehler: 3.478313e-06

```

```

stuexp3_xb_jorgensen <- getstuexp3(
  p = yb_jorgensen/100,
  q = xb_jorgensen,
  start = best_stuexp3_xb_jorgensen,
  show.output = TRUE,

```

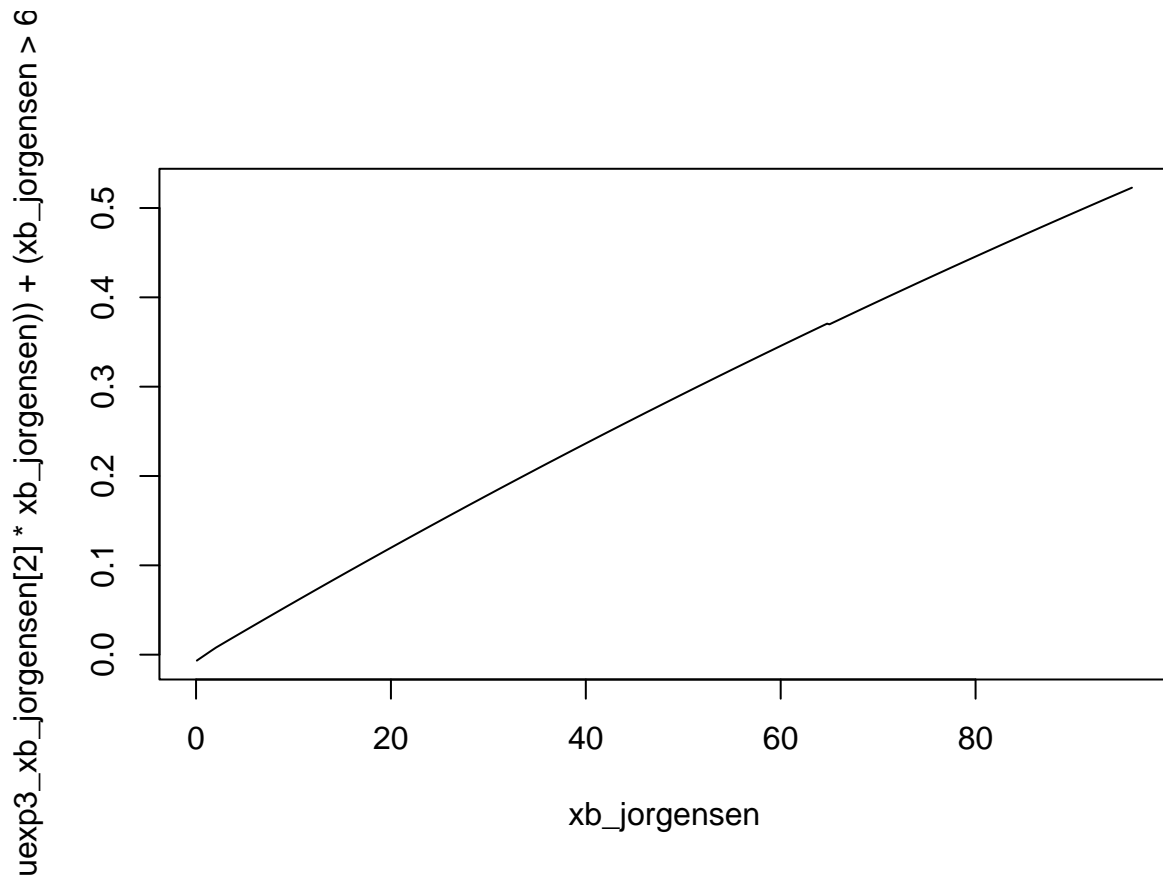
```
## $par
## [1] 0.001000000 0.003955892 0.002505107
##
## $value
## [1] 3.478313e-06
##
## $counts
## function gradient
##      50      50
##
## $convergence
## [1] 52
##
## $message
## [1] "ERROR: ABNORMAL_TERMINATION_IN_LNSRCH"
```

```
##      1.para      2.para      3.para
## 0.001000000 0.003955892 0.002505107
```

```

plot(xb_jorgensen,
      ((xb_jorgensen > 0 & xb_jorgensen <= 2 ) * (1 - exp(-stuexp3_xb_jorgensen[1] * xb_jorgensen)) +
        (xb_jorgensen > 2 & xb_jorgensen <= 6 ) *
          (1 - exp(-stuexp3_xb_jorgensen[1] * 2)) +
          (exp(-2 * stuexp3_xb_jorgensen[2]) - exp(-stuexp3_xb_jorgensen[2] *
            xb_jorgensen)) +
          (xb_jorgensen > 6 & xb_jorgensen <= 65 ) *
            (1 - exp(-stuexp3_xb_jorgensen[1] * 2)) +
            (exp(-2 * stuexp3_xb_jorgensen[2]) - exp(-6 * stuexp3_xb_jorgensen[2] *
              xb_jorgensen)) +
            (exp(-6 * stuexp3_xb_jorgensen[3]) -
              exp(-stuexp3_xb_jorgensen[3] * xb_jorgensen))),
      type = "l")

```



## 2-Stufige Exponentialverteilung

```

# 2-Stufige Exponentialverteilung
source("C:/Users/Chau/Documents/Masterarbeit/R Funktionen/getstuexp2.R")

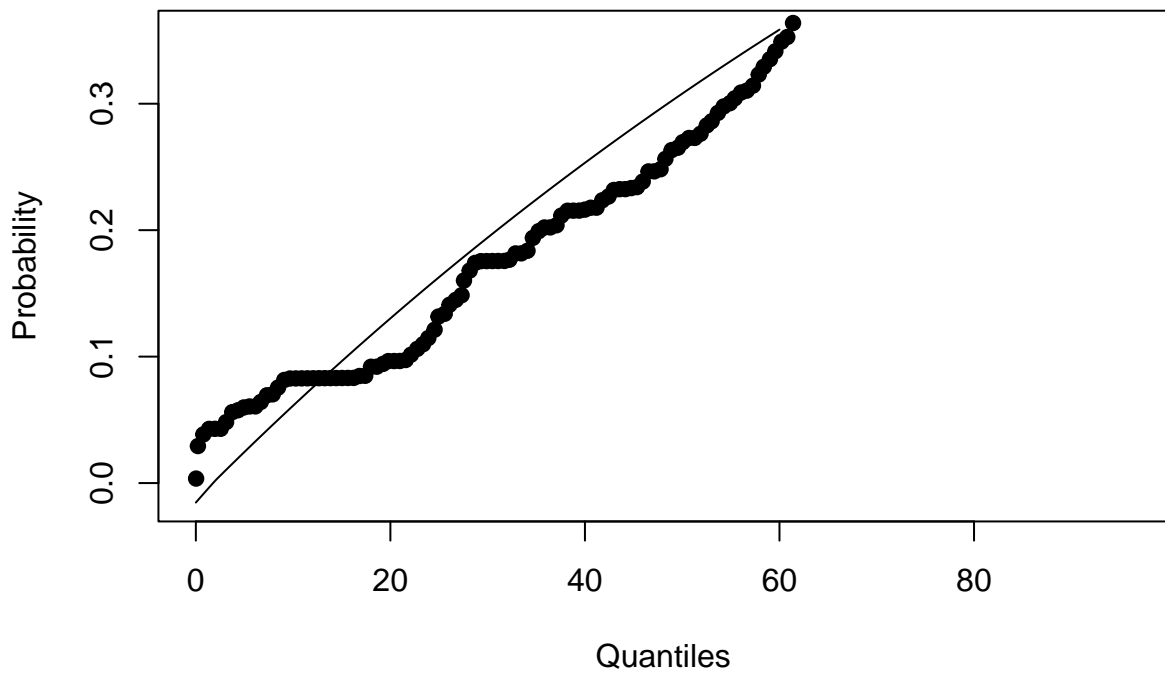
best_stuexp2_xr_jorgensen <- find_best_start_2parameter(p = yr_jorgensen/100,
  q = xr_jorgensen,
  max_beta = 1,
  max_eta = 0.5,
  steps_beta = 0.1,
  steps_eta = 0.01,
  fitting_function = getstuexp2)

```

```
stuexp2_xr_jorgensen <- getstuexp2(
  p = yr_jorgensen/100,
  q = xr_jorgensen,
  start = best_stuexp2_xr_jorgensen,
  show.output = TRUE,
  plot = TRUE,
  wert1 = 2
)
```

```
## $par
## [1] 0.001000000 0.007752908
##
## $value
## [1] 5.619892e-06
##
## $counts
## function gradient
##      8      8
##
## $convergence
## [1] 0
##
## $message
## [1] "CONVERGENCE: REL_REDUCTION_OF_F <= FACTR*EPSMCH"
```

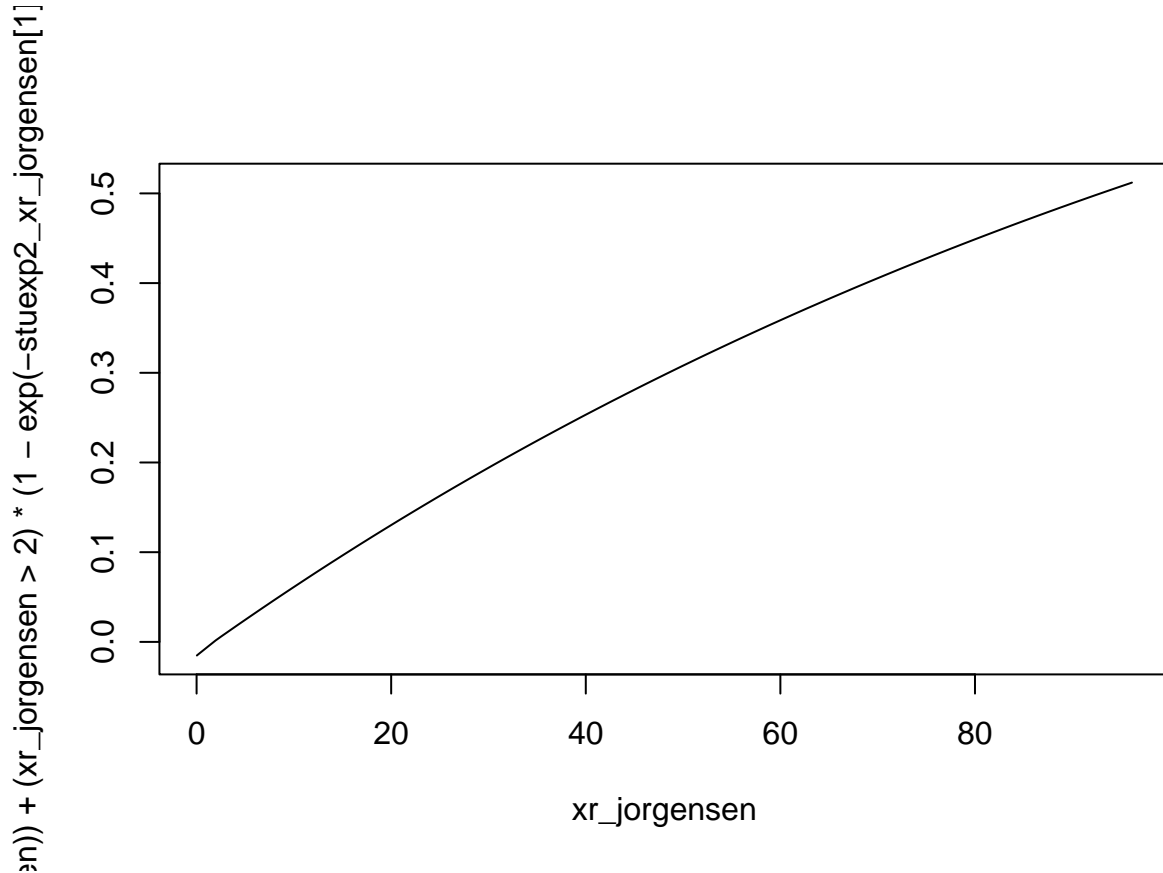
**2 stueckw. Exponential (1.para = 0.001, 2.para = 0.00775)**

[illegible]

```
stuexp2_xr_jorgensen
```

```
##      1.para      2.para  
## 0.001000000 0.007752908
```

```
plot(xr_jorgensen,  
      ((xr_jorgensen > 0 & xr_jorgensen <= 2) * (1 - exp(-stuexp2_xr_jorgensen[1] * xr_jorgensen)) +  
        (xr_jorgensen > 2) * (1 - exp(-stuexp2_xr_jorgensen[1] * 2)) +  
        (exp(-2 * stuexp2_xr_jorgensen[2]) -  
         exp(-stuexp2_xr_jorgensen[2] * xr_jorgensen))),  
      type = "l")
```



```
best_stuexp2_xb_jorgensen <- find_best_start_2parameter(p = yb_jorgensen/100,  
                                                         q = xb_jorgensen,  
                                                         max_beta = 1,  
                                                         max_eta = 0.5,  
                                                         steps_beta = 0.1,  
                                                         steps_eta = 0.01,  
                                                         fitting_function = getstuexp2)
```

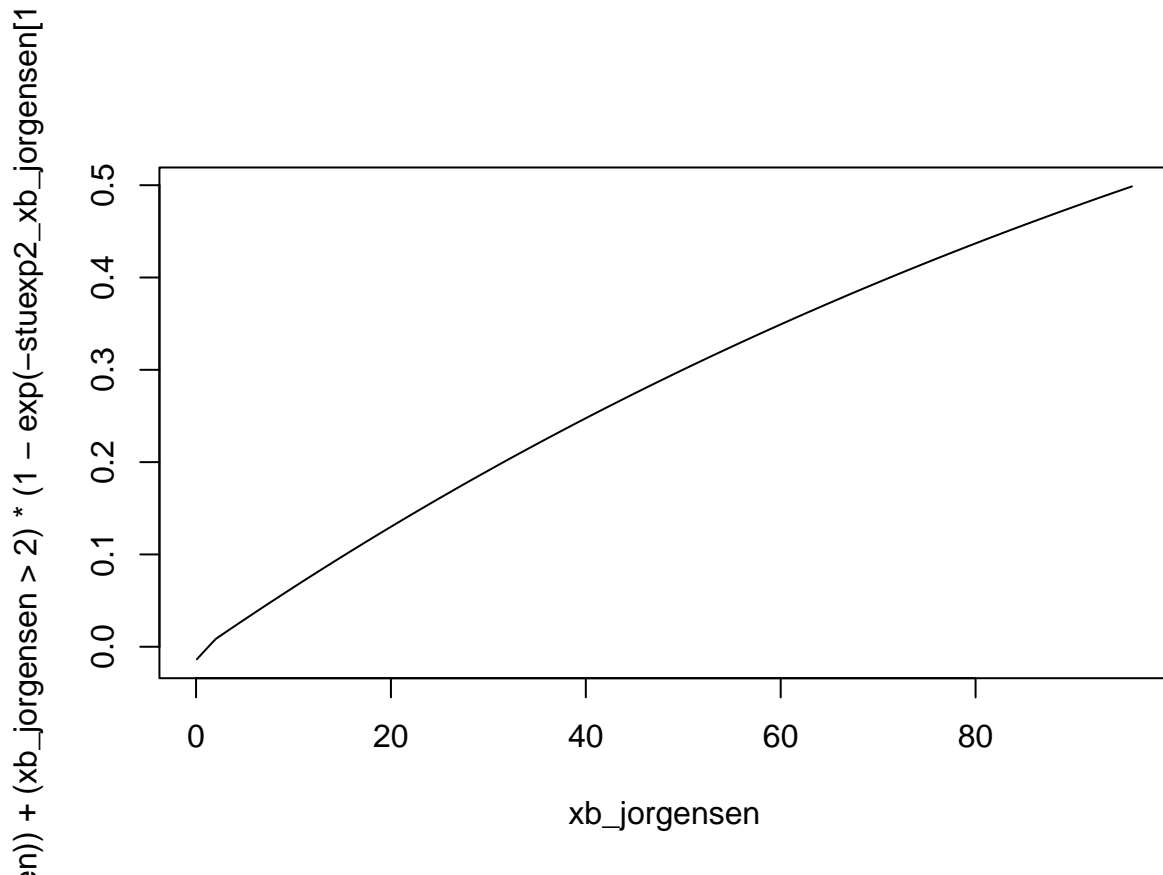
```
## Bester Startparameter: 0.8 0  
## Bester Fehler: 5.537071e-06
```

```
stuexp2_xb_jorgensen <- getstuexp2(  
  p = yb_jorgensen/100,  
  q = xb_jorgensen,  
  start = best_stuexp2_xb_jorgensen,
```



```
## $par
## [1] 0.004172411 0.007319418
##
## $value
## [1] 5.537071e-06
##
## $counts
## function gradient
##      28      28
##
## $convergence
## [1] 0
##
## $message
## [1] "CONVERGENCE: REL_REDUCTION_OF_F <= FACTR*EPSMCH"
```

```
plot(xb_jorgensen,
      ((xb_jorgensen > 0 & xb_jorgensen <= 2) * (1 - exp(-stuexp2_xb_jorgensen[1] * xb_jorgensen)) +
        (xb_jorgensen > 2) * (1 - exp(-stuexp2_xb_jorgensen[1] * 2)) +
        (exp(-2 * stuexp2_xb_jorgensen[2]) -
          exp(-stuexp2_xb_jorgensen[2] * xb_jorgensen))),
      type = "l")
```



## Mischung aus 2 Exponentialverteilungen

```
# Mischung aus 2 Exponentialverteilungen
source("C:/Users/Chau/Documents/Masterarbeit/R Funktionen/getexex.R")

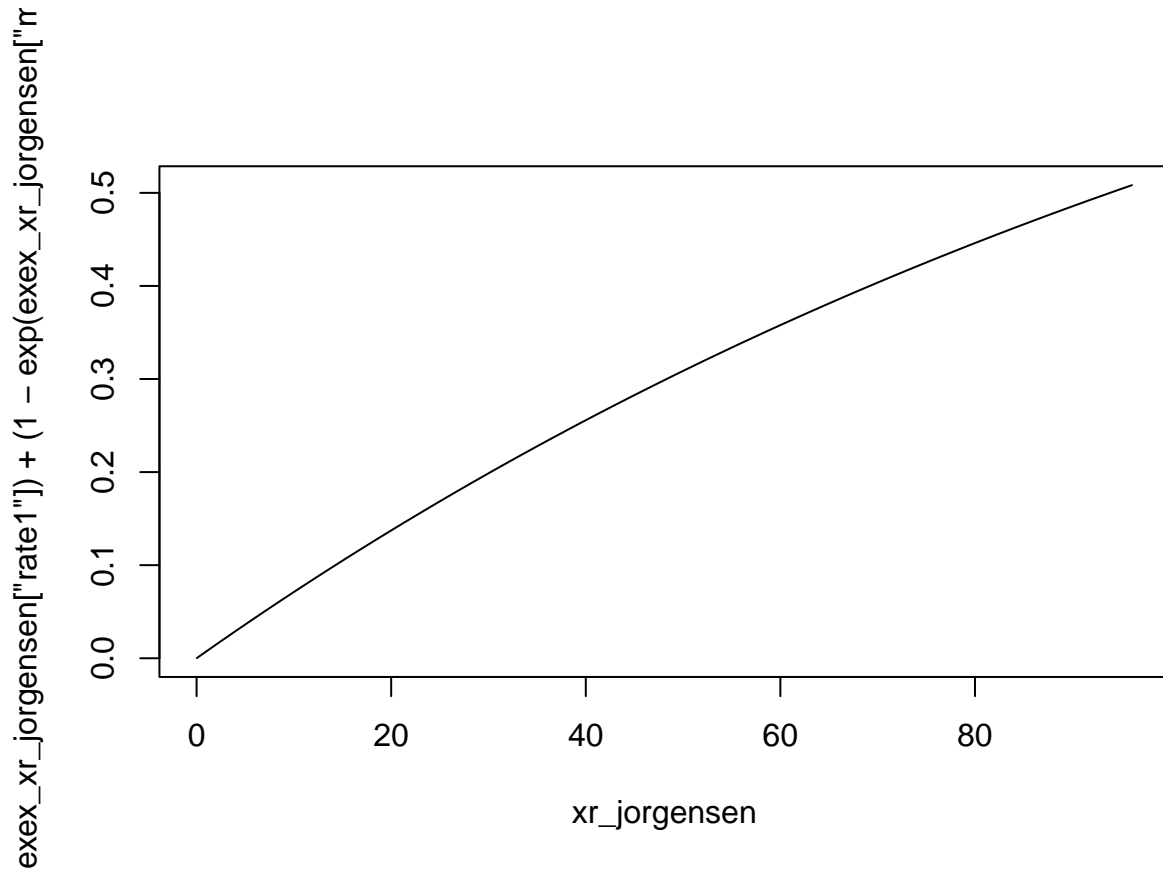
best_exex_xr_jorgensen <- find_best_start_3parameter(p = yr_jorgensen/100,
  q = xr_jorgensen,
  max_shape1 = 0.7,
  max_shape2 = 0.7,
  max_scale = 1,
  steps_shape1 = 0.1,
  steps_shape2 = 0.1,
  steps_scale = 0.1,
  fitting_function = getexex)

## Bester Startparameter: 0.6 0.3 0
## Bester Fehler: 6.120317e-06
```



```
## 0.007377371 0.007389922 0.001196593
```

```
plot(xr_jorgensen,
      (exp(exex_xr_jorgensen["mix"]) / ( 1 + exp(exex_xr_jorgensen["mix"]))) * stats::pexp(q = xr_jorgensen["rate"],
                                                rate = exex_xr_jorgensen["rate2"])),
      (1 - exp(exex_xr_jorgensen["mix"]) / ( 1 + exp(exex_xr_jorgensen["mix"]))) *
      stats::pexp(q = xr_jorgensen,
                  rate = exex_xr_jorgensen["rate2"])),
      type = "l")
```



```
best_exex_xb_jorgensen <- find_best_start_3parameter(p = yb_jorgensen/100,
                                                    q = xb_jorgensen,
                                                    max_shape1 = 0.7,
                                                    max_shape2 = 0.7,
                                                    max_scale = 1,
                                                    steps_shape1 = 0.1,
                                                    steps_shape2 = 0.1,
                                                    steps_scale = 0.1,
                                                    fitting_function = getexex)
```

```
## Bester Startparameter: 0.4 0.6 0.4
```

```
## Bester Fehler: 5.456246e-06
```

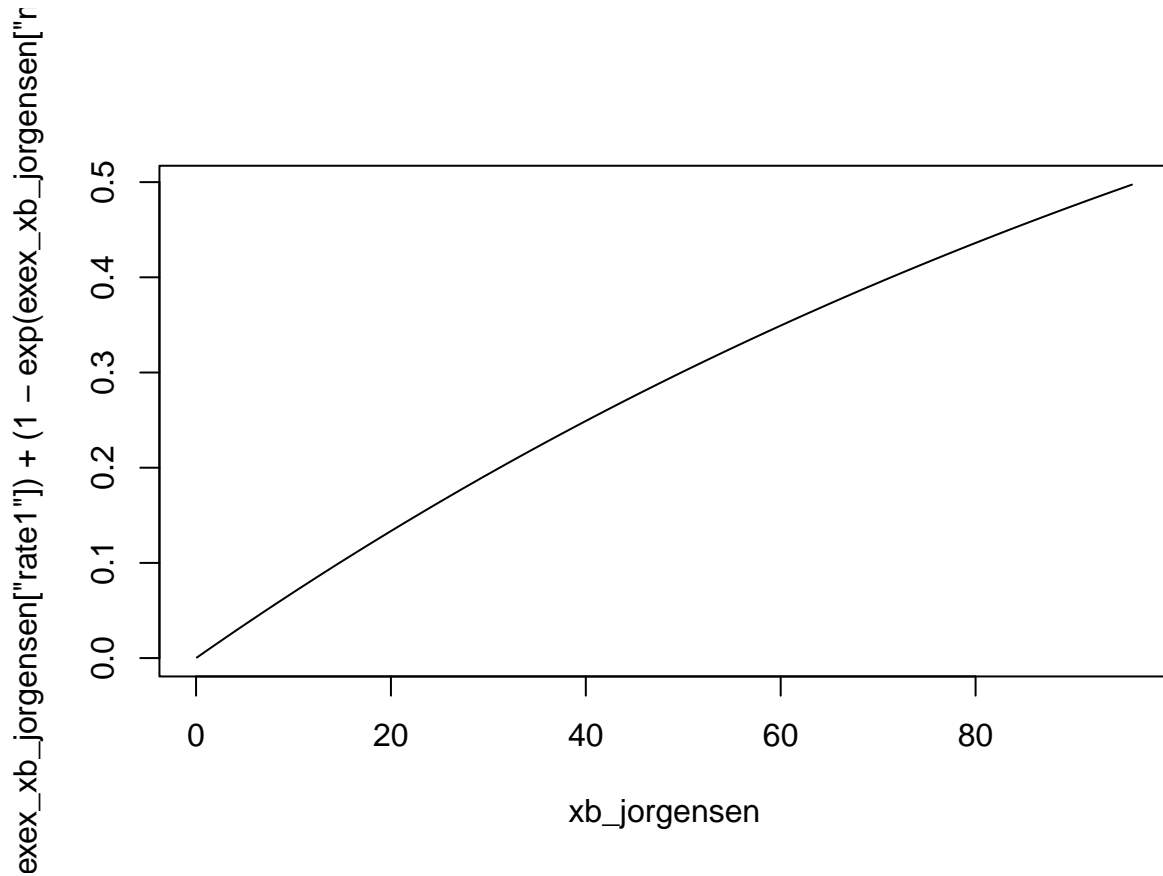
```
exex_xb_jorgensen <- getexex(
  p = yb_jorgensen/100,
  q = xb_jorgensen,
  start = best_exex_xb_jorgensen,
  show.output = TRUE,
```



```

      (1 - exp(exex_xb_jorgensen["mix"])) / ( 1 + exp(exex_xb_jorgensen["mix"]))) *
stats::pexp(q = xb_jorgensen,
           rate = exex_xb_jorgensen["rate2"])),
type = "l")

```



## Ergebnnis

```

getvalue <- function(p, q, best_start, fitting_function){
  if(identical(as.character(substitute(fitting_function)), "getstuexp2")){
    output <- capture.output({
      result <- getstuexp2(p = p, q = q, start = best_start, show.output = TRUE,
                          plot = FALSE, wert1 = 2)
    })
  }
  else if(identical(as.character(substitute(fitting_function)), "getstuexp3")){
    output <- capture.output({
      result <- getstuexp3(
        p = p, q = q, start = best_start,
        show.output = TRUE, plot = FALSE, wert1 = 2, wert2 = 6)
    })
  }
  else{
    output <- capture.output({
      result <- fitting_function(p = p, q = q, start = best_start,

```

```

        show.output = TRUE, plot = FALSE)
    })
}

# Berechne den Fehler (jorgensen_r_1$value) für die aktuellen Startparameter
error <- as.numeric(gsub("\\[1\\]\\s+", "", output[5]))

return(error)
}

best_test <- function(p, q, weibull, weib, weibex, weibex2, expweib, stuexp3, stuexp2,
                      exex, start_weibull, start_weib, start_weibex, start_weibex2,
                      start_expweib, start_stuexp3, start_stuexp2, start_exex,
                      group){
  weibull_val <- getvalue(p, q, start_weibull, getweibullpar)
  weib_val <- getvalue(p, q, start_weib, getweibpar)
  weibex_val <- getvalue(p, q, start_weibex, getweibex)
  weibex2_val <- getvalue(p, q, start_weibex2, get2weibex)
  expweib_val <- getvalue(p, q, start_expweib, getexpweib)
  stuexp3_val <- getvalue(p, q, start_stuexp3, getstuexp3)
  stuexp2_val <- getvalue(p, q, start_stuexp2, getstuexp2)
  exex_val <- getvalue(p, q, start_exex, getexex)

  error_distribution_pairs <- list(
    list(weibull_val, "W"),
    list(weib_val, "e.W."),
    list(weibex_val, "M. W&E"),
    list(weibex2_val, "M. e.W&E"),
    list(expweib_val, "e.W o. lambda = 1"),
    list(stuexp3_val, "3 s.E."),
    list(stuexp2_val, "2 s.E."),
    list(exex_val, "M. E&E")
  )

  # Suchen Verteilung mit dem kleinsten Fehler
  best_pair <- error_distribution_pairs[[which.min(sapply(
    error_distribution_pairs, function(pair) pair[[1]]))]]

  # Drucken Sie die Ergebnisse
  cat("Beste Verteilung:", best_pair[[2]], "\n")
  cat("Bester Fehler:", best_pair[[1]], "\n")
  cat("Gruppe: ", group)

  return(c(group, best_pair[[2]], best_pair[[1]]))
}

best_savr <- best_test(yb_jorgensen/100, xb_jorgensen, jorgensen_b_1, weibull_xb_jorgensen,
                      weibex_xb_jorgensen, weibex2_xb_jorgensen, expweib_xb_jorgensen,
                      stuexp3_xb_jorgensen, stuexp2_xb_jorgensen, exex_xb_jorgensen,
                      best_jorgensen_b_1, best_weibull_xb_jorgensen,
                      best_weibex_xb_jorgensen, best_weibex2_xb_jorgensen,
                      best_expweib_xb_jorgensen, best_stuexp3_xb_jorgensen,
                      best_stuexp2_xb_jorgensen, best_exex_xb_jorgensen, "SAVR")

```

```

## Beste Verteilung: M. W&E
## Bester Fehler: 5.537916e-07
## Gruppe: SAVR

best_tavr <- best_test(yr_jorgensen/100, xr_jorgensen, jorgensen_r_1, weibull_xr_jorgensen,
  weibex_xr_jorgensen, weibex2_xr_jorgensen, expweib_xr_jorgensen,
  stuexp3_xr_jorgensen, stuexp2_xr_jorgensen, exex_xr_jorgensen,
  best_jorgensen_r_1, best_weibull_xr_jorgensen,
  best_weibex_xr_jorgensen, best_weibex2_xr_jorgensen,
  best_expweib_xr_jorgensen, best_stuexp3_xr_jorgensen,
  best_stuexp2_xr_jorgensen, best_exex_xr_jorgensen, "TAVR")

## Beste Verteilung: M. e.W&E
## Bester Fehler: 8.776434e-07
## Gruppe: TAVR

tab <- matrix(c("NOTION", "alle", "TSM", best_tavr[1], best_tavr[2],
  best_tavr[3], weibex2_xr_jorgensen[1:3], NA, NA, NA,
  weibex2_xr_jorgensen[4],
  "NOTION", "alle", "TSM", best_tavr[1], "M. W&E",
  getvalue(yr_jorgensen/100, xr_jorgensen, best_weibex_xr_jorgensen, getweibex),
  NA, NA, weibex_xr_jorgensen[1:2], NA, weibex_xr_jorgensen[3:4],
  "NOTION", "alle", "TSM", best_savr[1], best_savr[2],
  best_savr[3], NA, NA, weibex_xb_jorgensen[1:2], NA,
  weibex_xb_jorgensen[3:4]),
  ncol=13, byrow=TRUE)

rownames(tab) <- NULL
colnames(tab) <- c('Studie', 'PG', 'EP', 'GR', 'Verteilung', 'SSE', '$\\alpha$',
  '$\\theta$', '$\\lambda_1$', '$\\lambda_2$', '$\\lambda_3$',
  '$\\vartheta$', '$\\psi$')

results <- as.data.frame(tab)

# Speichern
write.table(results, "results_jorgensen.txt", sep = "\t", row.names = FALSE)

# Funktion zur Überprüfung von NA-Werten für Zeichenketten und numerische Werte
is_non_empty <- function(x) {
  return(!is.na(x) & x != "")
}

# Spalten mit mindestens einem nicht-NA-Wert ermitteln
nicht_leere_spalten <- colSums(apply(results, is_non_empty)) > 0

# Konvertieren Sie die Tabelle in eine Markdown-Tabelle
print(results[, nicht_leere_spalten])

##   Studie   PG   EP   GR Verteilung      SSE      $\\alpha$
## 1 NOTION alle TSM TAVR   M. e.W&E 8.776434e-07 0.318085638347565
## 2 NOTION alle TSM TAVR     M. W&E 9.427333e-07          <NA>
## 3 NOTION alle TSM SAVR     M. W&E 5.537916e-07          <NA>
##      $\\theta$      $\\lambda_1$      $\\lambda_2$      $\\vartheta$
## 1 47.5324992810653 0.313042930144028          <NA>          <NA>
## 2              <NA> 1.55865720109492 112.219015159449 3.78019047741798

```



```
## 3          <NA> 1.6887982737076 116.055116397409 1.34734657059735
##          $\\psi$
## 1 2.43780761004889
## 2 3.02502395742585
## 3 2.56340038363206
```